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Please amend the specification under the provisions of 37 C.F.R. §1.121(b)(1) as follows.

On page 7, lines 3-15, please delete the paragraph which begins "Figure 1 . . ." and insert the following paragraph:

B1 -Figure 1 Nucleotide (SEQ ID NO: 1) and Deduced Amino Acid Sequence (SEQ ID NO: 2) of Dorsalin-1

The numbering of the protein sequence starts with the first methionine of the long open reading frame. The putative signal sequence is typed in bold letters. The RSKR (SEQ. ID No. 17) sequence preceding the proteolytic cleavage site (arrow) is underlined. The site of insertion of the 10 amino acid c-myc epitope is marked with an asterisk. The accession number for dorsalin-1 is L12032. --

On page 7, lines 17-35, please delete the paragraph which begins "Figure 2 . . ." and insert the following paragraph:

B2 -Figure 2 Dorsalin-1 is a Member of the TGF- β Superfamily

(A) Alignment of the COOH-terminal amino acid sequences of dorsalin-1 and some representative members of the TGF- β superfamily. Residues that are identical in at least 4 of the 7 proteins are printed in white on a black background. The 7 conserved cysteine residues are marked with an asterisk. Gaps introduced to optimize the alignment are represented by dashes. Known proteolytic cleavage sites in these proteins are marked with an arrow head. Numbers at the right indicate the number of amino acids present in the